

KOS Diagnostic Lab (A Unit of KOS Healthcare)



Dr. Vinay Chopra
MD (Pathology & Microbiology)
Chairman & Consultant Pathologist

Dr. Yugam Chopra MD (Pathology) CEO & Consultant Pathologist

		LABORATO	RY REPORT			
Name	:Mr RAJAT		Sex/Age	: Male/35 Years	Case ID	:40321603550
Ref By	:DR. VINAY CHO	PRA	Dis.Loc.	1	Pt ID	
Bill, Loc.	KOS DIAGNOS	TIC LAB			Pt. Loc.	3
Registratio	on Date & Time	: 23-Mar-2024 09:59	Sample Typ	e : Heparin Whole Blood - N	a Ph#	3
Sample Da	ite & Time	: 23-Mar-2024 09:59	Sample Co	II.By :	Ref Id	12
Report Da	te & Time	: 30-Mar-2024 12:59	Acc. Rema	rks :	Ref ld 2	4

Chromosome Analysis Report

Clinical History	No clinical history available.					
Karyotype (ISCN Nomenclature 2020)		46,XY				
Interpretation	Normal Karyotype					
Banding Method	: GTG	Culture Type	: 72hrs PHA stimulated			
Banding Method Banding Resolution	: GTG : Approx 550	Culture Type Metaphases Counted	: 72hrs PHA stimulated			
			: 72hrs PHA stimulated : 20 : 05			

For specimens received from non NCGM locations, it is presumed that it belongsto the patient as identified on the labels of the container/Test Requisition Formand it has been verified as per GCLP (Good Clinical Lab Practices) by the referrer at the time of collection of the specimen. NCGM's responsibility is limited to the analytical part of the assay performed.

Dr. Sandip Shah M.D. (Path, & Bact.) Consultant Pathologist

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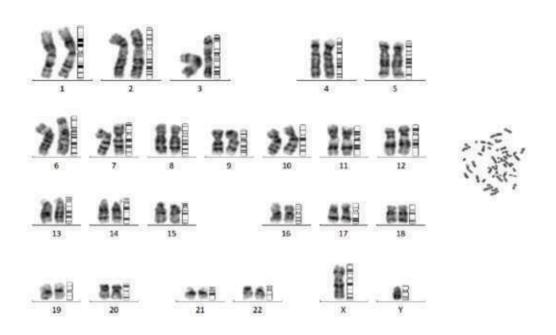


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Ref By	DR. VINAY CHO	OPRA	Dis.Loc.	3	Pt ID		
Bill. Loc.	:KOS DIAGNOST	TIC LAB			Pt. Loc.		
Registration	Date & Time	: 23-Mar-2024 09:59	Sample Typ	e : Heparin Whole Blood - N	a Ph#	3	
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Karyogram and Metaphase



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Limitation

The error rate of the test is 0.5%. The normal report does not rule out very Low grade mosaicism, minor chromosomal anomalies, and deletion, Duplication or Inversion at very subtle level. The report should be interpreted in accordance with the counselling provided before the test and with the report. A standard G-banded Karyotype usually has a resolution of around 5 Mb.

Disclaimer

Polymorphic variants have not been reported as these variants are not associated with specific disease or phenotype. Cytogenetically visible polymorphic variants include variants involving heterochromatin (variant size), satellite size, pericentric inversions (heterochromatic or euchromatic regions) [e.g., 1qh+/qh-, 9qh+/qh-, 16qh+/qh-, acrocentric p+ or p-, Yqh+/qh-, inv(9)(p11q13); inv(2)(p11.2q13)] and also euchromatic variants (e.g., located on 4p16, 8p23.1, 9p12, 9q13-q21.12, 15q11.2, 16p11.2).

Reference: Silva, M., de Leeuw, N., Mann, K., Schuring-Blom, H., Morgan, S., Giardino, D., Rack, K. and Hastings, R., 2019. European guidelines for constitutional cytogenomic analysis. European Journal of Human Genetics, 27(1), pp 1-16.

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-- End Of Report -----

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